

Ultrastructure and molecular biological changes of chronic gastritis, gastric cancer and gastric precancerous lesions: a comparative study

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Abstract

AIM: To carry out a comparative study on ultrastructure and molecular biological changes of chronic gastritis (CG), gastric cancer (GC) and gastric precancerous lesions.

METHODS: By the use of histochemical staining, SEM with EDAX, TEM with EDAX, image analysis technique, RIA and chemiluminescence method, gastric mucosa of 168 patients were synchronously analyzed in morphology, trace elements, DNA, cAMP, SOD, ³H-TdR LCT and serum LPO were also done.

RESULTS: The incidence of epithelial nucleoplasmic ratio >1, lobulated nuclei, inter-chromatin aggregation of granules, nucleolar hypertrophy, and the content of DNA, Zn, Cu in nuclei and serum LPO of each group were showed as follows: normal control group (0.0, 0.0, 6.7, 0.0, 12.6±2.7, 7.6±0.4, 58.4±0.3, 2.6±0.6), CSG group (5.7, 2.9, 7.4, 2.9, 15.2±3.1, 8.1±0.5, 58.9±0.5, 4.2±0.7), CAG group (31.3, 29.7, 45.3, 42.2, 16.5±3.1, 8.6±0.4, 59.3±0.5, 4.5±0.6), CA group (100.0, 100.0, 72.2, 50.0, 30.7±8.2, 8.8±0.3, 59.5±0.4, 6.8±1.6), ATP⁺⁺ group (61.5, 38.5, 23.1, 38.5, 23.5±8.9, 8.3±0.4, 59.1±0.4, 5.1±1.2), IM⁺⁺⁺ ATP⁺⁺ group (77.8, 55.5, 33.3, 44.4, 25.1±7.2, 8.4±0.5, 59.5±0.4, 6.5±1.1), IM⁺⁺⁺ ATP⁺⁺ group (100.0, 100.0, 75.0, 62.5, 28.5±9.1, 8.9±0.5, 59.7±0.4, 7.6±0.7), IMII_b group (100.0, 62.5, 75.0, 50.0, 27.3±10.3, 8.6±0.3, 59.5±0.4, 6.1±0.9); whereas the content of Zn, Cu in mitochondria and cAMP, SOD in gastric mucosa, and ³H-TdR LCT of each group were shown as follows: normal control group (9.2±0.5, 58.3±0.3, 15.9±1.5, 170.5±6.1, 1079.7±227.4), CSG group (8.6±0.5, 57.8±0.3, 14.6±1.8, 163.3±5.6, 867.3±240.5), CAG group (8.3±0.4, 57.5±0.3, 13.4±1.8, 161.2±4.3, 800.9±221.8), CA group (8.9±0.4, 57.1±0.3, 10.2±3.9, 152.2±3.8, 325.7±186.8), ATP⁺⁺ group (9.1±0.4, 57.0±0.3, 12.4±1.8, 161.5±3.8, 642.9±174.3), IM⁺⁺⁺ ATP⁺⁺ group (8.6±0.4, 56.9±0.3, 12.0±2.3, 152.2±2.5, 326.3±160.3), IM⁺⁺⁺ ATP⁺⁺ group (8.5±0.3, 56.8±0.2, 10.4±0.9, 147.4±2.6, 316.1±170.7), IMII_b group (8.6±0.3, 56.9±0.3, 11.9±1.9, 150.0±2.8, 318.9±145.8), there were significant differences between groups ($P < 0.05-0.01$).

CONCLUSION: There was a significant difference between CG and GC in their ultrastructure and molecular biology. Only on the condition of changes of internal environment in combination with the harmful effect of external

environment, chronic atrophic gastritis can then develop into gastric cancer. Hence it might have similar epithelial cell ultrastructure and molecular biological changes in ATP⁺⁺, IMII_b and cancer, hence there were similar patterns of occurrence, development and transformation. Recognition of this trend might help to explore problems of prevention and cure.

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INTRODUCTION

We conducted histochemical staining, detection of cAMP, SOD and DNA of gastric mucosa of 168 patients with chronic gastritis (CG) or gastric cancer (GC); SEM and TEM with EDAX were used to synchronously determine ultrastructures and trace elements and image analysis technique was used to determine nuclear DNA. ³H-TdR LCT and serum LPO were also carried out. Following was the comparative study.

MATERIALS AND METHODS

Materials

168 patients definitely diagnosed as suffering from CSG, CAG and GC were our subjects of study. According to "The standards for the classification of chronic gastritis, for gastrofiberscopic diagnosis and for histopathological diagnosis of atrophic gastritis", gastric mucosa tissue slices underwent HE staining and AB_{PH2.5}/PAS, Hid/AB_{PH2.5} and HiD/AB_{PH2.5} histochemical staining. ATP was classified into low-grade, middle-grade and heavy-grade and IM into IM I_a, IM I_b, IM II_a, IM II_b. 68 cases were definitely diagnosed as CSG, including 43 males, 25 females, average age 43 years, average course of disease 4a, 26 cases without IM or ATP; 31 cases with concomitant or solitary IM-19 cases low-grade, 10 cases middle-grade and 2 cases heavy-grade; 9 cases with solitary ATP-8 cases low-grade, 1 case middle-grade; 2 cases with IM and concomitant ATP, IM and ATP were both middle-grade. 64 cases were definitely diagnosed as CAG, including 37 males, 27 females, average age 47 years average course of disease 6a, 3 cases without IM or ATP; 39 cases with concomitant solitary IM (17 cases low-grade, 4 cases middle-grade and 18 cases heavy-grade); 8 cases with solitary ATP (2 cases low-grade, 6 cases middle-grade.) 14 cases with IM concomitant ATP (12 cases middle-grade IM, 2 cases heavy-grade IM, 8 cases low-grade ATP, 6 cases middle-grade ATP). 36 cases were definitely diagnosed as GC, including 22 males, 14 females, average age 52 years, average course of disease, 2a, all with IM or ATP. 12 cases with solitary IM (6 cases low-grade, 3 cases middle-grade, 3 cases heavy-grade); 7 cases with solitary ATP (1 case low-grade, 6 cases middle-grade);

17 cases with IM concomitant ATP (5 cases middle-grade IM, 12 cases heavy-grade IM; 7 cases low-grade ATP, 9 cases middle-grade ATP, 1 case heavy-grade ATP). Of the 45 healthy volunteers who underwent gastroscopy and biopsy, 15 cases had basically normal gastric mucosa tissues, including 6 males 9 females, average age 37, and they were referred to as normal control (NC) group.

Methods

Under gastroscopy, three pieces of gastric mucosa were taken from the focal and nonfocal areas in antral region and body of stomach as specimens for section preparation, SEM, TEM, detection of DNA, cAMP and SOD. Blood was taken for ^3H -TdRLCT and LPO. For the observation of gastric mucosa ultrastructure and determination of its trace elements^[1-4], 501B SEM with 9100/60 EDAX was used to observe the three pieces of specimens of each patient, under the direct vision of SEM, the EDAX probe automatically detected the samples within 0.1-0.01 mm² range all the elements under 12 in atomic number and automatically calculated the weight percentage (WT%) of each element in element series. 15 points were fixed in the three pieces of mucosa of every patient to carry out 15 detections, and the average WT% of each element was taken as its actual WT%. In every detection, 21 elements such as Na, Mg, Al, Si, P, S, K, Ca, Fe, Cu, Zn, Ti, Cr etc were detected, and weight percentage (WT%) of each element between elements of gastric mucosa was calculated.

By using EM430 TEM with 9100/60 EDAX, three pieces of mucosa specimens of every patient were magnified in unison by five magnifying powers (3 600, 7 200, 14 000, 19 000, 29 000) to randomly take the pictures of the panoramagram, local area and organelle and to detect the atomic number percentage (AT%) of each nuclear and mitochondrial element between trace elements^[5-10]. After gastric mucosa cell smear was stained by Feulgen staining, IBAS 2000 image analysis technique was adopted to detect IOD, which is taken as the relative content of nuclear DNA^[7,8]. RIA was adopted to detect the content of gastric mucosa cAMP (Pmol/g); chemiluminescence method was adopted to detect the activity of SOD (u/g)^[4]. ^3H -TdRLCT. (Bq/L whole blood) was carried out; thiobarbituric acid development process was adopted to detect serum LPO (u mol/L)^[4].

Statistical method

χ^2 and t test.

RESULTS

Histopathology of gastric mucosa

Between NC group. CSG group. CAG group and CA group, there were significant differences in the incidence rate and degree of different types of IM and ATP in glands propria of gastric mucosa ($P < 0.05$ -0.001, Table 1). The surface of normal gastric mucosa was isolated by crisscross small groves into many lesser gastric areas, assuming convolution shape (Figure 1). In these areas, there were many gastric pits (the mouths of gastric glands), which were shaped like craters. The concave walls had round or oval epithelial cells of almost the same size. In CSG gastric mucosa there were scattered denatured, diabrotic and necrotic exfoliated epithelial cells, on whose surface S-shape *Helicobacter pylori* (HP) were found (Figure 2). Massive epithelial cells became diabrotic, anabrotic and exfoliated forming micro ulcers. The ulcers spread out from their centers, with adjacent cells crushed, destructed in irregular shapes and arrangement. In CSG gastric mucosa, cyst was found accidentally (Figure 3), the epithelial cells of crater walls were atrophic and denatured,

of different sizes and deranged; the cells became diabrotic and necrotic and had inflammatory cell infiltration and the glands propria of the serious cases were shaped like grid framework structure (Figure 4). The surfaces of the epithelial calls of IM gastric mucosa were thickly coated, villi were invisible and the intercellular boundaries were not clear (Figure 5). SEM was not able to definitely indentify gastric mucosa ATP, but only able to distinguish whether cells were hyperplastic or not. Degeneration, hyperplasia or cellular regeneration occurred in gastric mucosa. The hyperplastic cells are of different sizes and states and the active memberanes were shaped like lesser tubercles (Figure 6, 7). The cells of gastric cancer were different sizes and shapes, deranged, characterized by obvious heteromorphic nature, conglobated and shaped like grapes or cauliflower. There were nearby or basal infiltration, destruction and ulceration of cancer cells, which have developed into ulcers.

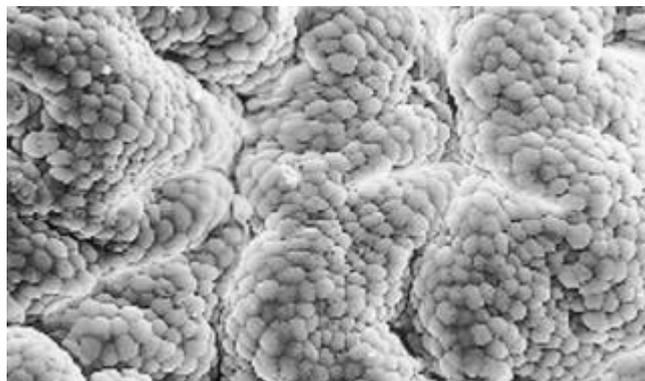


Figure 1 Convolution shape of mucosa surface. $\times 320$.

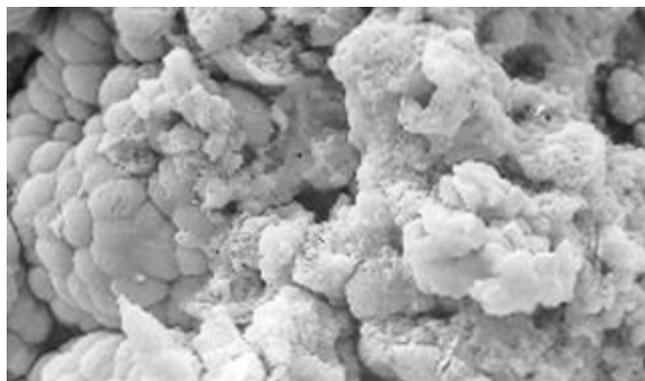


Figure 2 Degeneration, diabrosis, necrosis and exfoliation of gastric mucosa epithelial cells. $\times 640$.

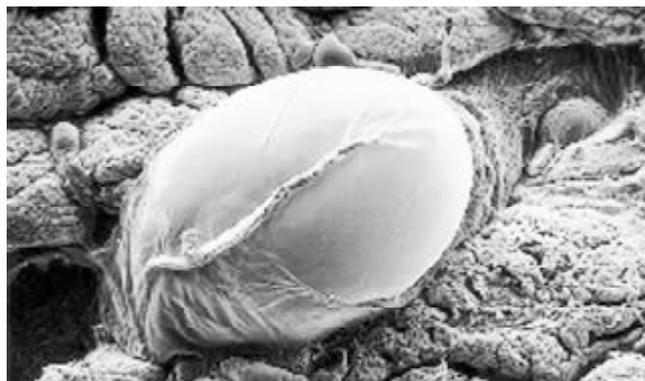


Figure 3 Cysts of gastric mucosa. $\times 64$.

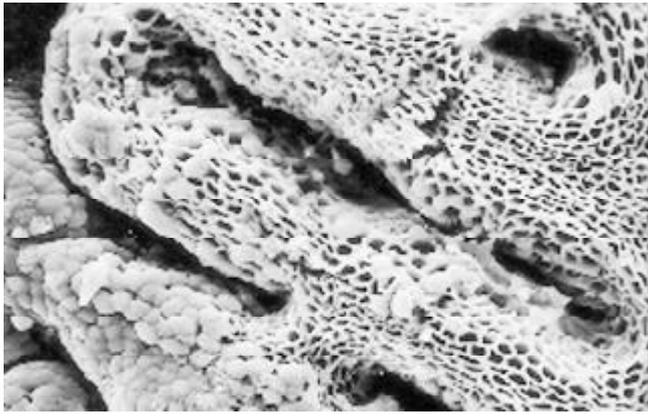


Figure 4 Atrophic and denatured glands propria in grid framework structure. $\times 320$.

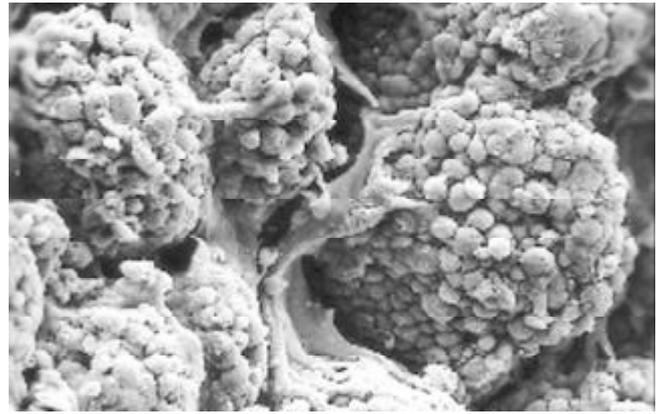


Figure 8 The cells of gastric cancer are conglomerated and shaped like grapes. $\times 320$.

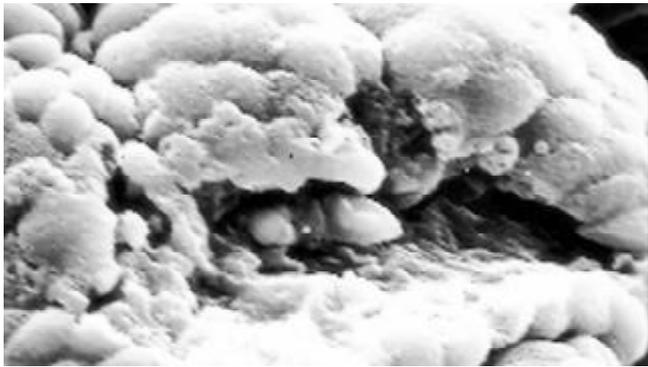


Figure 5 The surface of IM cells is thickly coated, intercellular boundaries are not clear. $\times 1\ 250$.

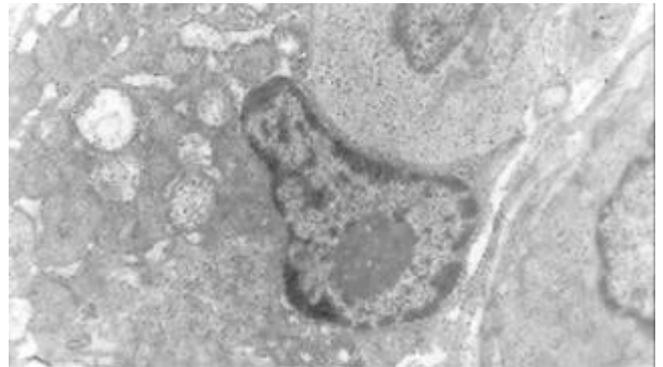


Figure 9 There is an increase in inter chromatinic granules densification, nucleolar granules become thick nucleoli expand with irregular margins. $\times 14\ 000$.

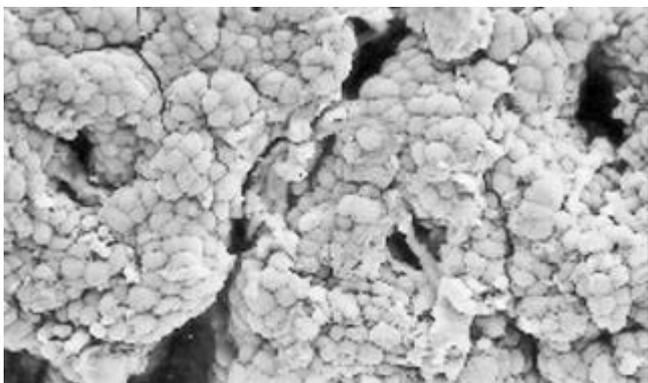


Figure 6 The hyperplastic cells are of different sizes and states, the active membranes are shaped like lesser tubercles. $\times 320$.

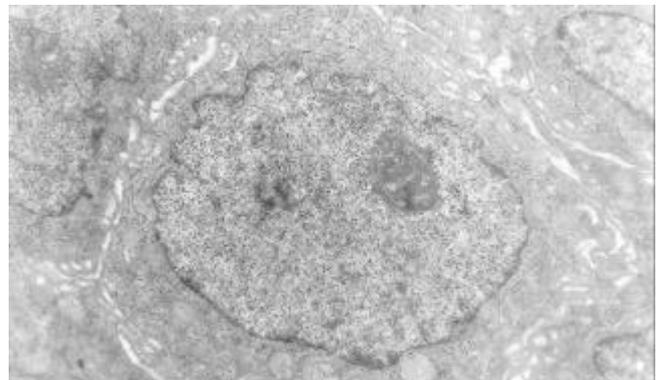


Figure 10 Nucleolar margination. $\times 7\ 200$.

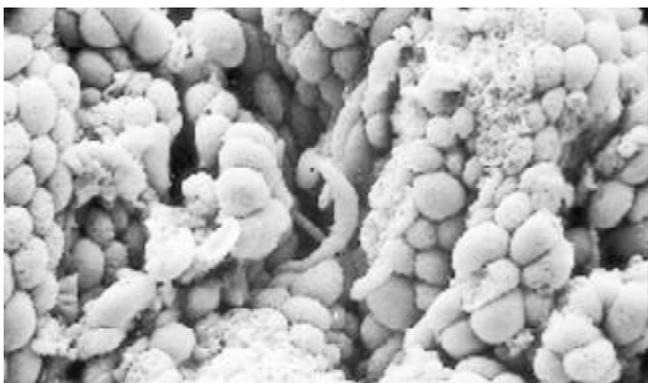


Figure 7 The hyperplastic cells are of different sizes and states, the active membranes are shaped like lesser tubercles. $\times 640$.



Figure 11 Multi nucleoli. $\times 14\ 000$.

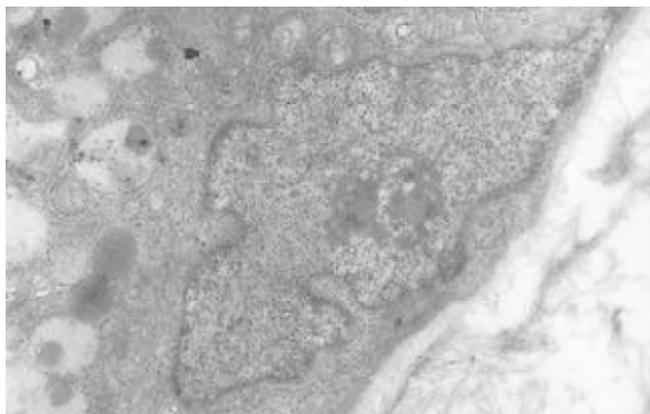


Figure 12 Nucleolar division. $\times 14\ 000$.

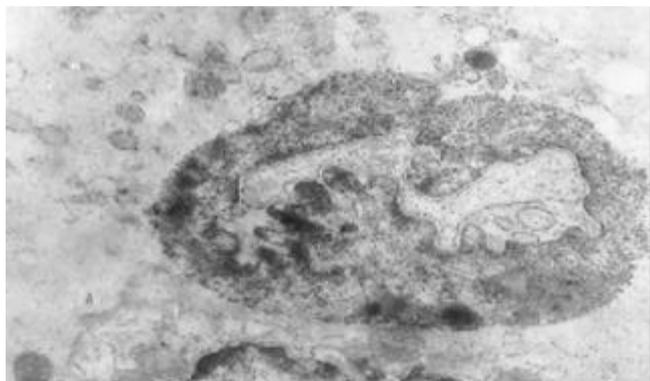


Figure 13 Intranuclear inclusion. $\times 14\ 000$.

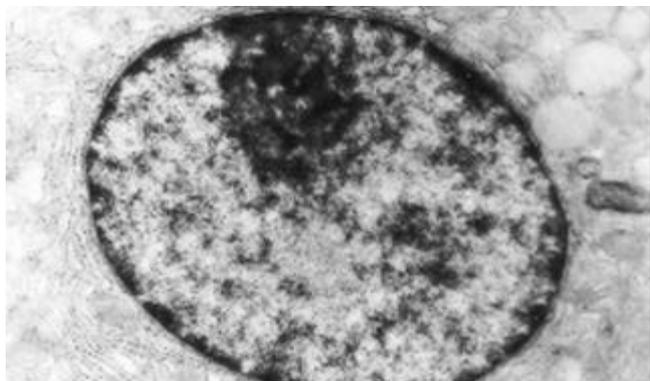


Figure 14 The densification of inter chromatinic grandules. $\times 14\ 000$.

The ultrastructure of gastric mucosa epithelial cells

The mucous cells on epithelial cell surface of relatively normal gastric mucosa were columnar epithelial cells covering endogastric surfaces and inside wall of gastric pit. Their free surfaces had short micro villi; the mucous neck cells were distributed over the necks of gastric glands, and on the tops of these cells were a few short thick micro villi. Chief cells were distributed over the bodies and bottoms of gastric glands; parietal cells were big and conic and their conical tops turned towards gland cavities; the endocrine cells lied between chief cells and parietal cells and they were small, and their nuclei are round or oval, the nuclei envelopes are slightly bent, lobulated nuclei were few, the nucleocytoplasmic ratio was less than 1.

The nuclear chromatins were scattered, or associated with nucleoli or along nuclear perimeters. The light bright zones between heterochromatins in the nucleoli are euchromatins; the nucleoli had high electron density without capsule.

Mitochondria were round or oval, scattering around nuclei. The mitochondria consisted of outer membrane, inner membrane, outer ventricle, inner ventricle and cristae. Crista, the inward folded inner membrane, was a hollow canal leading to outer ventricle. Some mitochondrial cristae directly led to cytoplasm. Cristae were generally in tabular arrangement, parallel to each other and vertical to mitochondrial long axes. In cytoplasm were many parallel rough surfaced endoplasmic reticulus (RERs), secretory granules, well-developed Golgi's bodies and scattered mitochondria. The free surfaces of epithelial cells of CG gastric mucosa had dropped micro villi; the intercellular space expanded and cell conjunctions decreased; the karyoplasmic ratio of both ill-differentiated epithelial cells and IM cells was greater than 1, and lobulated nuclei increased; nucleolar granules became thick and nucleoli expanded with irregular margins (Figure 9).

There occurred nucleolar margination (Figure 10), multinucleoli (Figure 11) and nucleolar division (Figure 12). The obsolescent epithelial nuclei shrank and the karyoplasmic ratio was still less; the heterochromatins lied densely around nuclei; the electron density was low in the center of nuclei and nuclei were loop in shape. (referred to as chromatin margination); the shrunk nuclei were denticular, in which the electron density was moderately homogeneous and chromatins were not found (referred to as chromatin homogenization). There were mitochondrial swelling, hypertrophy, pyknosis, hyaline degeneration as well as vacuolar degeneration. The deformed mitochondrias were in C-shape or U-shape. There were zigzag, longitudinal, sparse and pyknosis cristae, and deranged cristas. There was an decrease in the number of mitochondrias and their cristae. There were mitochondrial decreases in number, mitochondrial swelling, or crista fragmentation, vacuolar degeneration, and RERs in circular arrangement; the Golgi's bodies become atrophic and had lost their typical structures; the cytoplasmic incretory granules decreased. There were still greater epithelial cell changes in cancer cells of gastric cancer than in cells of gastric mucosa of chronic gastritis, intranuclear inclusion appeared (Figure 13). There was an increase in peri-chromatinic and interchromatinic granular densification (Figure 14). In euchromatins; there was a significant difference between CSG group and CAG group in nucleolar margination and nucleolar looping ($P < 0.05-0.001$, Table 2, 3); between IM II_b group, ATP⁺⁺ group, IM⁺⁺+ATP⁺⁺ group and IM⁺⁺⁺+ATP⁺⁺ group, compared with NC group, CSG group and CAG group, the difference was also significant ($P < 0.05-0.001$, Table 2, 3). There is no significant difference between IM⁺⁺⁺+ATP⁺⁺ group, IM II_b group and CA group ($P > 0.05$, Table 2, 3).

The biochemical assay in gastric mucosa and serum

The content of nuclear Zn and Cu increased progressively in the sequence of NC group, CSG group, CAG group and CA group; the content of mitochondrial Zn and Cu decreased progressively in the same sequence. There was significant differences between these groups ($P < 0.05-0.001$, Table 4). The content of Zn, Cu, cAMP and SOD in gastric mucosa decreased progressively in the same sequence; the content of DNA increased progressively in the above sequence. There were significant differences between these groups ($P < 0.05-0.001$, Table 5). The content of ³H-TdRLCT decreased in the sequence of NC group, CSG group, CAG group and CA group; content of serum LPO increased in the same sequence; there was significant difference between these groups ($P < 0.05-0.001$, Table 6). Between IMII_b group, ATP⁺⁺ group, IM⁺⁺+ATP⁺⁺ group and IM⁺⁺⁺+ATP⁺⁺ group, NC group, CSG group and CAG group, there was also significant difference ($P < 0.05-0.001$, Table 4, 5, 6). Between IM⁺⁺⁺+ATP⁺⁺ group, IM II_b group and CA group, There was no significant difference ($P > 0.05$, Table 4, 5, 6).

Table 1 Comparison between classification of IM and grading of ATP in CG and GC gastric mucosa *n* (%)

Groups	<i>n</i>	IM classification				Grading of ATP						
		I _a	I _b	II _a	II _b	ATP ⁺	ATP ⁺⁺	IM ⁺⁺⁺ +ATP ⁺	IM ⁺⁺⁺ +ATP ⁺⁺	IM ⁺⁺⁺ +ATP ⁺	IM ⁺⁺⁺ +ATP ⁺⁺	IM ⁺⁺⁺ +ATP ⁺⁺⁺
NC	15	2(13.3)	1(6.7)									
CSG	68	16(23.5)	8(11.8)	6(8.8)	3(4.4)	8(11.8)	1(1.5)		2(2.9)			
CAG	64	16(25.0) ^b	11(17.2) ^b	13(20.3) ^d	13(20.3) ^d	2(3.1) ^d	6(9.4) ^d	7(10.9)	5(7.8) ^c	1(1.5)	1(1.6)	
CA	36		4(11.1)	9(25.0) ^c	16(44.4) ^{df}	1(2.8) ^d	6(16.7) ^d	3(8.3)	2(5.6) ^c	4(11.1) ^f	7(19.4) ^f	1(2.8)

Either Solitary IM or IM with concomitant ATP in CSG, CAG or CA is included in IM classification statistically, compared with NC group. ^a*P*<0.05, ^b*P*<0.01; compared with CSG group. ^c*P*<0.05, ^d*P*<0.01; compared with CAG group. ^e*P*<0.05, ^f*P*<0.01. In table 2-6, the marks are the same as here.

Table 2 The ultrastructures of epithelial nuclei of gastric mucosa in CG and GC *n* (%)

Group	<i>n</i>	Appearance		Chromatin		Nucleons	
		Nucleoplasmic ratio >1	Nucleus lobulated	Margination or homogeneity	Perinuclear concentrated	Hypertrophy or margination	looping
NC	15			1(6.7)	1(6.7)		
CSG	68	5(5.7)	2(2.9)	7(10.3)	5(7.4)	2(2.9)	2(2.9)
CAG	64	20(1.3) ^d	19(9.7) ^d	33(51.6) ^{bd}	29(45.3) ^{bd}	27(42.2) ^d	14(21.9)
CA	36	36(100.0) ^{bf}	36(100.0) ^{bf}	17(47.2) ^d	26(72.2) ^{bdf}	18(50.0) ^d	17(47.2) ^{df}
IMII _b	8	8(100.0) ^b	5(62.5) ^h	5(62.5) ^b	6(75.0) ^b	4(50.0)	3(37.5)
ATP ⁺⁺	13	8(61.5) ^{hi}	5(38.5) ^{hj}	4(30.8) ^{bgi}	3(23.1) ^{bhj}	5(38.5)	1(7.7) ^{hj}
IM ⁺⁺⁺ +ATP ⁺⁺	9	7(77.8) ^{hhik}	5(55.5) ^{hk}	3(33.3) ^{bgi}	3(33.3) ^{bhj}	4(44.4)	3(33.3) ^l
IM ⁺⁺⁺ +ATP ⁺⁺	8	8(100.0) ⁱⁿ	8(100.0) ^{ikm}	5(62.5) ^{bkm}	6(75.0) ^{blm}	5(62.5) ^{lm}	3(37.5) ^l

Compared with CA group, ^g*P*<0.05, ^h*P*<0.01, Compared with IMII_b group, ⁱ*P*<0.05, ^j*P*<0.01, Compared with ATP⁺⁺ group, ^k*P*<0.05, ^l*P*<0.01. Compared with IM⁺⁺⁺+ATP⁺⁺ group, ^m*P*<0.05, ⁿ*P*<0.01. In table 3-6, the marks are the same as here.

Table 3 Mitochondria ultrastructures of gastric mucosa epithelial cells in CG and GC ($\bar{x}\pm s$)

Group	<i>n</i>	Number	Swelling or Ovegrowth (%)	Matrix fading (%)	Vacuolar degeneration (%)	Pyknosis (%)	Crista number	Fragmentation and derangement of cristas(%)
NC	15	86.5±27.3	3.4±1.6	3.0±1.1	2.9±1.9	1.1±0.8	12.8±3.2	2.2±1.1
CSG	68	65.4±21.1 ^b	7.2±3.8 ^b	7.8±5.0 ^b	6.4±4.5 ^b	1.9±0.9 ^b	8.2±3.2 ^b	5.8±3.1 ^b
CAG	64	52.2±20.8 ^{bd}	10.9±4.5 ^{bd}	11.7±8.6 ^{bd}	11.6±7.7 ^{bd}	3.7±1.1 ^{bd}	6.9±3.5 ^{bc}	8.9±3.7 ^{bd}
CA	36	38.8±31.5 ^{bde}	12.9±4.2 ^{bde}	15.6±5.4 ^{bdf}	14.6±5.8 ^{bde}	3.9±0.6 ^{bd}	8.1±1.9 ^{bde}	10.5±2.7 ^{bde}
IMII _b	8	36.8±30.8 ^b	13.8±3.2 ^b	14.1±4.3 ^b	12.5±3.9 ^b	3.3±0.8 ^{bg}	9.3±2.1 ^b	7.6±1.1 ^{bh}
ATP ⁺⁺	13	49.1±27.9 ^b	11.3±2.9 ^b	8.3±3.5 ^{bh}	9.3±4.7 ^b	1.9±0.5 ^{bhj}	11.3±1.8 ^{hi}	5.2±1.2 ^{bhj}
IM ⁺⁺⁺ +ATP ⁺⁺	9	41.3±30.8 ^b	12.8±2.1 ^b	12.1±2.8 ^{bg}	11.2±3.7 ^b	2.3±0.3 ^{bhjk}	11.0±1.5 ^{ahi}	6.2±1.3 ^{bhi}
IM ⁺⁺⁺ +ATP ⁺⁺	8	35.4±31.5 ^b	13.3±4.3 ^b	15.2±5.3 ^{bi}	12.8±4.3 ^b	3.9±0.5 ^{bbin}	8.4±2.1 ^{bin}	8.6±1.0 ^{bhin}

Table 5 Gastric mucosa Zn, Cu, DNA, cAMP and SOD in CG and GC ($\bar{x}\pm s$)

Group	<i>n</i>	Zn (WT%)	Cu(WT%)	DNA(IOD)	cAMP(pmol/g)	SOD(u/g)
NC	15	4.1±1.0	5.2±0.8	12.6±2.7	15.9±1.5	170.5±6.1
CSG	68	2.8±1.9 ^b	4.0±1.5 ^a	15.2±3.1 ^b	14.6±1.8 ^b	163.3±5.6 ^b
CAG	64	2.0±1.8 ^{bc}	3.4±1.5 ^{bd}	16.5±3.1 ^{bc}	13.4±1.8 ^{bc}	161.2±4.3 ^{bc}
CA	36	1.5±1.2 ^{bde}	2.8±0.9 ^{bde}	30.7±8.2 ^{bdf}	10.2±3.9 ^{bdf}	152.2±3.8 ^{bdf}
IMII _b	8	1.7±0.9 ^b	3.5±0.9 ^b	27.3±10.3 ^b	11.9±1.9 ^b	150.0±2.8 ^b
TP ⁺⁺	13	2.7±1.3 ^{bhi}	4.4±0.9 ^{ahi}	23.5±8.9 ^{bg}	12.4±1.8 ^{bg}	161.5±3.8 ^{aj}
IM ⁺⁺⁺ +ATP ⁺⁺	9	2.3±1.5 ^b	4.1±0.9 ^{bh}	25.1±7.2 ^b	12.0±2.3 ^b	152.2±2.5 ^{bl}
IM ⁺⁺⁺ +ATP ⁺⁺	8	1.5±0.8 ^{bkm}	2.9±0.8 ^{bhin}	28.5±9.1 ^b	10.4±0.8 ^{bil}	147.4±2.6 ^{bhin}

Table 4 Nuclear and mitochondrial Zn, Cu of gastric mucosa in CG and GC ($\bar{x}\pm s$)

Group	n	Nuclear		Mitochondria	
		Zn (AT%)	Cu(AT%)	Zn (AT%)	Cu(AT%)
NC	15	7.6±0.4	58.4±0.3	9.2±0.5	58.3±0.3
CSG	68	8.1±0.5 ^b	58.9±0.5 ^b	8.6±0.5 ^b	57.8±0.3 ^b
CAG	64	8.6±0.4 ^{bd}	59.3±0.5 ^{bd}	8.3±0.4 ^{bd}	57.5±0.3 ^{bd}
CA	36	8.8±0.3 ^{bde}	59.5±0.4 ^{bde}	8.9±0.4 ^{bde}	57.1±0.3 ^{bde}
IMII _b	8	8.6±0.3 ^b	59.5±0.4 ^b	8.6±0.3 ^b	56.9±0.3 ^b
ATP ⁺⁺	13	8.3±0.4 ^{bhi}	59.1±0.4 ^{bhi}	9.1±0.4 ⁱ	57.0±0.3 ^b
IM ⁺⁺ +ATP ⁺⁺	9	8.4±0.5 ^{bg}	59.5±0.4 ^{bk}	8.6±0.4 ^{bi}	56.9±0.3 ^b
IM ⁺⁺⁺ +ATP ⁺⁺	8	8.9±0.5 ^{bl}	59.7±0.4 ^b	8.5±0.3 ^{agl}	56.8±0.2 ^{bh}

Table 6 LPO and ³H-TdRLCT of CG and GC ($\bar{x}\pm s$)

Group	n	LPO(umol/L)	³ H-TdRLCT(Bq/L)
NC	15	2.6±0.6	1079.7±227.4
CSG	68	4.2±0.7 ^b	867.3±240.5 ^b
CAG	64	4.5±0.6 ^{bc}	800.9±221.8 ^{bc}
CA	36	6.8±1.6 ^{bdf}	325.7±186.8 ^{bdf}
IMII _b	8	6.1±0.9 ^b	318.9±145.8 ^b
ATP ⁺⁺	13	5.1±1.2 ^{bhi}	642.9±174.3 ^{bhi}
IM ⁺⁺ +ATP ⁺⁺	9	6.5±1.1 ^b	326.3±160.6 ^{bk}
IM ⁺⁺⁺ +ATP ⁺⁺	8	7.6±0.7 ^{bgilm}	316.1±170.7 ^{bl}

DISCUSSION

Chronic gastritis, CAG in particular, is generally acknowledged as precancerous state. However, not all CAGs are likely to develop into gastric cancer because there is significant difference between chronic gastritis and gastric cancer in the ultrastructure and molecular biology of gastric mucosa. In the grading and classification of IM, grading of ATP of gastric mucosa, incidence rate of epithelial karyoplasmic ratio >1, incidence rate of perichromatinic granular densification and of nucleolar hypertrophy, the number of mitochondria and cristae as well as the incidence rate of mitochondrial degeneration; as well as in the quantitative changes of nuclear DNA, Zn, Cu, cAMP, SOD and serum LPO and ³H-TdRLCT, there were significant differences between NC group, CSG group, CAG group, CA group, IM II_b group, ATP⁺⁺ group, IM⁺⁺+ATP⁺⁺ group and IM⁺⁺⁺+ATP⁺⁺ group ($P<0.05-0.0001$); There was no significant difference between IM⁺⁺⁺+ATP⁺⁺ group, IM II_b group and CA group ($P>0.05$). There are similar epithelial ultrastructures and molecular biochemical changes in ATP⁺⁺, IM II_b and CA. Consequently they should have similar law of occurrence, development and transformation, which might contribute to the exploration of prevention and treatment of the disease.

Clinically, only a few chronic gastritis patients' gastric mucosa epithelial cells are likely to develop into cancer cells, but it is essential that there should be a process which causes genetic variation of cells, the key of which is the quantitative changes of epithelial nuclear DNA. The DNA molecular is coded with nucleotide sequence fragments of special genetic codes, every fragment is a functional group, referred to as gene. The gene order has great stability. Only when intracellular cAMP declines, cytodifferentiation is disturbed. The metabolism of Zn, Cu in the body is disturbed, which in turn disturbs the enzyme system and inhibits the synthesis and activity of SOD; because of the NADPH oxidation reduction circulation and the catalytic function of xanthine oxidase, a great deal of oxygen free radicals are produced far beyond the

cleaning ability of SOD. The excessive accumulated oxygen free radicals react in lipid peroxidation with unsaturated fatty acid of inner and outer membranes of mitochondria, producing LPO. Thus level of serum LPO rises. The inner and outer membranes of mitochondria are destructed, causing decrease and derangement of mitochondrial cristae, mitochondrial degeneration decrease of adenosine triphosphate production, and inadequate energy supply of gastric mucosa epithelial cells. When nuclei take up more Zn and Cu, the protein synthesis is enhanced nuclear division and hyperplasia are accelerated; through influencing lymphocyte metabolism, the quantitative changes of Zn and cAMP in turn inhibit lymphocyte transformation and result in the decline of ³H-TdRLCT level and immunity of the organism. The above changes of intraorganism environment, only if stimulated by certain external causes such as ionizing radiation, chemical injuries, infection S-shape HP and virus, are likely to cause the change of sequences in gene group, which is referred to as gene mutation; and genetic codes change accordingly, the changed codes are transferred to heterogenous cell descendants, causing cellular differentiation disturbance and even cancerous change. Human body is an integrity organism, in which the components of tissue and cell, and the bioactive substances including trace elements, enzymes, hormones, immunity and messenger substances exist with a definite content and a definite quantitative ratio. It's important to measure the "absolute" content of each bioactive substance, but it's more important to measure quantitative ratio of each bioactive substance because it can explain the trends and laws of variation of the whole body and inner circumstance^[9,10]. These ratios in normal body keep in a state of dynamic balance within a definite range. If this balance is broken, pathological phenomena and functional disturbance will arise. The ratio fluctuation of bioactive substances higher or lower than normal value, can lead to a series of ratio variation, and then cause both a definite disease diagnosed by modern medicine and clinical expression of Chinese medical syndrome (a synthesis of symptoms without peculiarity). The composite nature of Chinese medicine treats disease through adjusting the above mentioned pathological ratio is one of it's therapeutic mechanisms. This is why Chinese medicines have multilevel, multitarget, two-way adjusting effects on human body, and how the medicines regulate pathological ratio. It's a slow process for Chinese medicines to regulate pathological ratio, so It's therapeutic effect is slow too. Measuring the "quantitative ratio" of a series of bioactive substances not only will be the research direction of medical science and biological science, but also the research kernel of the basic theory of traditional Chinese medicine^[11-13].

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